06-DEC-2002; 2002US-0431620P. 05-DEC-2003; 2003WO-GB005323. 24-JUN-2004.

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맑
                                                                                                                                                                                                                                                                                                                                                                        RESULT 10
ADR13967/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xiao
US2004121341-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 198 BP; 56 A; 36 C; 49.G; 57 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 16; 81pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptides that interact with myelin proteins Nogo, TNR and MAG, useful in preparing a composition for treating CNS damage, spinal cord injury or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-468811/44.
P-PSDB; ADQ16419.
                                                                                      CDS
                                                                                                                                                                                                                                                                    Human NOGO-66
                                                                                                                                                                                                                                                                                                23-SEP-2004
                                                                                                                                                                                                                                                                                                                              ADR13967;
                                                                                                                                                                                                                                                                                                                                                          ADR13967 standard; cDNA; 198 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   major myelin proteine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating central nervous system (CNS) damage, spinal cord injury or stroke. The peptides may also be used in vaccines against myelin antigens. The vaccine is based on the specific inhibitory portions of major myelin proteins, instead of the whole protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tenascin-R (TN-R) (specifically TN-R epidermal growth factor like (TNR-BGFL)) and myelin-associated glycoprotein (MAG). These proteins have neural growth inhibitory activity. The peptide is isolated from a 7-mer phage display library exposed to a plate coated with the target protein. Peptides of the invention are useful in preparing a composition for treating central nervous system (CNS) damage, spinal cord injury or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  describes peptides which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence encodes Nogo-66 domain b.
                                                                                                                                  Homo sapiens
                                                                                                                                                                             multiple sclerosis; Creutzfeldt-Jacob disease;
multiple system atrophy; Lou Gehrig's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SIGE-)
                                                                                                                                                              progressive
                                                                                                                                                                                                      neural regeneration; apoptosis; amylotrophic lateral sclerosis;
Alzheimer's disease; Parkinson's disease; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DENI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (specifical)
                                                                                                                                                                                                                                     gene; human; myelin-associated glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                    ű
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CUGGAUAGCUUGGAUCACACCCUUG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SINGAPORE GEN HOSPITAL PTE DENISON C M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ly the Nogo-66 domain), the extracel (TN-R) (specifically TN-R epidermal)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                               supranuclear
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                    CDNA.
                                                                                                   Location/Qualifiers
                             note= "No start and stop codons given"
                                                                        *tag=
                                             product= "NOGO-66"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interact with the myelin proteins Nogo
                                                                                                                                                               palsy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Score 25; DB Pred. No. 0.07 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 12; Length 198; .079;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification
                                                                                                                                                                                              Kuru;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            matrix glycoprotein
h factor like (TNR-
                                                                                                                                                                                                                          ; neural growth; sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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PPA PA XXX
  The invention relates to a myelin-associated glycoprotein (MAG)
CC derivative comprising a mutation in or flanking MAG Ig-like domain 5
CC (Igd5), excluding the MAG derivative MAG (dl-3)-Fc, where the mutation
CC reduces or eliminates the ability of the derivative to regulate neurite
CC outgrowth as compared to endogenous or soluble MAG without eliminating
CC promoting neural growth and regeneration. They are also useful for
CC promoting neural degeneration associated with injuries, disorders, or
CC diseases. The disorder, disease, or condition is associated with
CC apoptosis or results from a demyelinating disease and includes
CC amylotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease,
CC funtington's disease, multiple sclerosis, Creutzfeldt-Jacob disease,
CC kuru, multiple system atrophy, amylotrophic lateral sclerosis (Lou
CC Gehrig's disease), or progressive supranuclear palsy. The present
CC sequence represents the human NOGO-66 CDNA.
Sequence 198 BP; 56 A; 36 C; 49 G; 57 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             New myelin-associated glycoprotein (MAG) derivative comprises a mutation in or flanking MAG Ig-like domain 5 (Igd5), excluding the MAG derivative MAG (d1-3)-Fc, useful promoting neural growth and regeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Filbin MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-2002; 2002US-00327213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 10; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P-PSDB; ADR13968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-479666/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-2002; 2002US-00327213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CAOZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FILB/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILBIN M T.
DOMENICONI M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAO Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Domeniconi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cao
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밁 Ś Matches Query Match Best Local 3 18; Similarity | CUGGAUAGCUUGGAUCACACCCUUG 25 CTGGATAGCTTGGATCACACCCTTG 9 Conservative 100.0%; 7; Mismatches Score 25; ub ... No. 0.079; DB 12; 0; Length 198; Indels 0, Gaps

0

RESULT 11 AAV23697/c 24-JUL-1998 AAV23697 AAV23697; standard; cDNA; 261 BP (first entry)

Human NSPLP protein coding sequence fragment.

NSPLP; neuroendocrine-specific protein-like protein; human; neurodegenerative disease; amyotrophic lateral sclerosis; c gene therapy;

Homo sapiens.

WO9806841-A2

19-FEB-1998

24-JUL-1997; 97WO-US013469.

12-AUG-1996; 9605-00700607

(INCY-) INCYTE PHARM INC

WPI; 1998-159533/14.

Bandman O,

Au-Young

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Goli SK,

Hillman J;

Human neuro-endocrine-specific protein-like proteins - useful for diagnosis, monitoring and treatment of cancer and neuro-degenerat

and neuro-degenerative

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RESULT 12
AAX41193/c
ID AAX41193 s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NSPLP are used for diagnosis and monitoring treatment of diseases associated with NSPLP expression, in usual immunoassays, and to isolate NSPLP from natural sources. The NSPLP proteins, or their fragments can also be used in drug screening to identify NSPLP antagonists. The nuclei acid can be used diagnostically and for monitoring treatment (in hybridisation or amplification assays); to isolate closely related hybridisation or amplification assays); to isolate closely related sequences; in gene therapy for both sense and antisense applications sequence (including use of ribozymes) and for mapping the natural genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein (NSPLP) of the invention. Recombinant Cells transcribed when the NSPLP proteins, which are used to treat DNA are used to express the NSPLP proteins, which are used to treat cancer and neurodegenerative diseases such as amyotrophic lateral scancer and neurodegenerative diseases such as amyotrophic lateral scancer and neurodegenerative diseases such as amyotrophic lateral scancer and neurodegenerative nucleic acids and antagonists of NSPLP can scance and antagonists of MSPLP can use the number of diseases used to inhibit activity of the NSPLP protein treatment of diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 45; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 261
                                                                                                                                                                                                                                                                                                                                                       forensic; gene therapy; chromosome mapping; signal peptide; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopoiesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted protein 5' EST SEQ ID NO:137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-1999
                                                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; secreted protein;
                                                                                                                                                                                                                                                                                        WO9906548-A2
New nucleic acids encoding human secreted proteins - obtained from cDNA libraries prepared from e.g. liver, ovary, brain, prostate, kidney, lung umbilical cord, placents and colon tissue.
                                                                                                                                                                                                                                                          11-FEB-1999.
                                                                                      WPI; 1999-153778/13
                                                                                                                    Dumas Milne Edwards
                                                                                                                                                                                       01-AUG-1997;
                                                                                                                                                                                                                       31-JUL-1998;
                                                                      P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gequence encodes a human neuroendocrine-specific protein-like
sin (NSPLP) of the invention. Recombinant cells transformed with
                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTGGATAGCTTGGATCACACCCTTG 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cuggauageuuggaucacacccuug 25
                                                                        AAY12360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                         97US-00905135
                                                                                                                                                                                                                          98WO-IB001222.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA; 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 59 C; 56 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                         Duclert
                                                                                                                                                                                                                                                                                                                                                                                                                                                EST; expressed sequence tag; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and for mapping the natural genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                67 T; 0 U; 17 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25;
                                                                                                                             Lacroix B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The nucleic
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sequences can be used for producing secreted human gene products. The can also be used to develop products for diagnosis and therapy. The proteins obtained may have cytokine activity, cell proliferation/differentiation activity, haematopolesis regulating activity, tissue growth regulating activity, reproductive hormone regulating activity, themostatic chemokinetic activity, haemostatic regulating activity, receptor/ ligand activity, anti-inflammatory activity, tumour inhibition activity or other activities. The product
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 319; 824pp; English
                                            can be used in forensic, gene therapy and chromosome mapping procedures. The sequences can also be used for obtaining corresponding promoter sequences. The nucleic acids encoding the signal peptide can be used for directing extracellular secretion of a polypeptide or the insertion of a polypeptide into a membrane, or importing a polypeptide into a cell
                                                                                                                                                                                                                                                                                                                                                                                                  AAX41094 to AAX41347 represent 5' expressed sequence tags (ESTs) for human secreted proteins, and encode the proteins given in AAX12261 to AAX12514, respectively. The proteins given represent the signal peptide and an N-terminal fragment of a secreted protein. The nucleic acid.
           111 T; 0 U; 0 Other;
                                                                                                                                                                                                        The products
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밁 8 Best Loc Matches Query Match Best Local (347 18; μ Similarity CIGGATAGCITGGATCACACCCITG cuggauagcuuggaucacacccuug 25 Conservative 72.0%; 100.0%; Pred. Score Mismatches ŏ. 25; 0 멂 2 0 Length 404; Indels 0 Gaps

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Sequence 404 BP; 110 A; 75 C;

108 G;

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RESULT 13
AAF90323/c
ID AAF903
XX
neuromuscular disorder; psychiatric disorder; developmental disorder; neuroprotective; nootropic; neuroleptic; antiparkinsonian;
                                                                                                                                                                                                                                                 Human NOGO-C CDNA.
                                                                                                                                                                                                                           NOGO-C; human; chromosome 2p21; peuropathy; spinal injury; brain injury; stroke; neuronal degeneration; Alzheimer's disease; Parkinson's disease;
                                                                                                                                                                                                                                                                                  AAF90323;
                                                                                                                                                                                                                                                                                                  AAF90323
                                                                                                                                                                                                                                                                   23-JUL-2001
                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                     cerebroprotective;
                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                  CDNA; 600
                                                                                                                                                                                                      neuroleptic; diagnosis;
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WO200136631-A1. 15-NOV-1999; 99GB-00026995 24-JAN-2000; 2000GB-00001550 14-NOV-2000; 2000WO-GB004345. (SMIK) SMITHKLINE BEECHAM PLC 25-MAY-2001 99GB-00026995.

Michalovich D,

Prinjha

æ

2001-343822/36.

New polypeptide designated NOGO-C is a splice variant of the human NOGO gene and may be useful in the treatment of neural disorders including Alzheimer's and Parkinson's diseases. P-PSDB; AAB82348

Claim 1; Page 25; 25pp; English

The present sequence is that of cDNA encoding human NOGO-C (see

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and polynucleotides, and methods for producing such polypeptides by recombinant techniques. Also disclosed are methods for utilising NOGO-C polypeptides and polynucleotides in the treatment of diseases including neuropathies, spinal injury, brain injury, stroke, neuronal degeneration, for example Alzheimer's disease and Parkinson's disease, neuromuscular disorders, psychiatric disorders and developmental disorders. Also provided are methods for identifying agonists and agonists for use in treating conditions associated with NOGO-C imbalance, and diagnostic assays for detecting diseases associated with inappropriate NOGO-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB82348). NOGO-C
chromosome 2p21. 2
previously been id
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene #3485 used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               activity or levels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CUGGAUAGCUUGGAUCACACCCUUG 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTGGATAGCTTGGATCACACCCTTG 192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . 2 Other splice variants, NOGO-A and NOGO-B, have identified. The invention provides NOGO-C polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>1</u>:8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnose liver cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
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disease Gene; liver cancer; ds; hepatocellular carcinoma; hepatotropic; metastatic liver tumour; cytostatic; expression profile; diseas progression; tumour; cytostatic;
ion; drug toxicity; d drug expression profile; disease st drug efficacy; drug metabolism state;

Homo sapiens

WO200229103-A2

02-OCT-2001; 2001WO-US030589

02-OCT-2000; 2000US-0237054P

(GBNB-) GENE LOGIC

Alvares C, Peres-Da-Silva ά Vockley JG;

WPI; 2002-426119/45

Diagnosing and detecting the progression carcinoma or metastatic liver tumor in a level of expression of two or more genes of liver cancer, hepatocellular patient, involves detecting the in a liver tissue sample.

Claim 1; SEQ ID NO 3485; 298pp; English.

The invention relates to a novel method for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver tumour in a patient, and differentiating metastatic liver cancer from hepatocellular carcinoma in a patient, involving detecting the level of expression of two or more genes represented in ABN93503-ABN97455 in a tissue sample. The method of the invention has hepatotropic, and cytostatic activity. The method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to monitor disease states, disease progression, drug toxicity, drug efficacy and drug metabolism. Note: The sequence data

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RESULT 15
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18-MAY-2001; 2001WO-US016450 PEPSDB; 19-MAY-2000; 2000US-0205515P (HUMA-) HUMAN GENOME 2002-122018/16. ABB89192. ξ SCI INC

prevention of neural, immune system, muscular, reproductive,
gastrointestinal, pulmonary, cardiovascular, renal and proliferative Novel 1405 isolated polypeptides, useful for diagnosis, treatment disorders.

Claim 4; SEQ ID NO 163; 2081pp + Sequence Listing; English.

The invention relates to novel genes (ABL89449-ABL90853) and proteins (CRB889040-ABB90444) useful for preventing, treating or ambliorating comedical conditions e.g. by protein or gene therapy. The genes are comedical conditions e.g. by protein or gene therapy. The genes are comedicated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and covarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune chisorders e.g. Addison's disease, allergies, autoimmune haemolytic canaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, canditiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing infectious diseases such as viral, bacterial, fungal and parasitic confections. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly comedicated in the printed specification, but was obtained in electronic format directly at ftp.wipo.int/pub/published_pct_sequences